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US-10-665- 611 VSLKRLRIPLSHEELEDPSIERRPVKDGCGGNTSITVRNATPTWARSDDPTLNGITFSIPEG  
US-10-665- 611 VSLKRLRIPLSHEELEDPSIERRPVKDGCGGNTSITVRNATPTWARSDDPTLNGITFSIPEG  
US-10-665- 611 VSLKRLRIPLSHEELEDPSIERRPVKDGCGGNTSITVRNATPTWARSDDPTLNGITFSIPEG  
consensus  
US-10-665- 672 ALVAVGVGCGKSSLLSALLAEMDKVEGHVAIKGSVAVYVPOQAWIONDSIRENILFGCOL  
US-10-665- 672 ALVAVGVGCGKSSLLSALLAEMDKVEGHVAIKGSVAVYVPOQAWIONDSIRENILFGCOL  
US-10-665- 672 ALVAVGVGCGKSSLLSALLAEMDKVEGHVAIKGSVAVYVPOQAWIONDSIRENILFGCOL  
US-10-665- 672 ALVAVGVGCGKSSLLSALLAEMDKVEGHVAIKGSVAVYVPOQAWIONDSIRENILFGCOL  
consensus  
US-10-665- 733 EEPYRSVIQACALLPDLIILPSGDRTEIGKGNLGGQKQKORVSLARAVYSNADIYLFDD  
US-10-665- 733 EEPYRSVIQACALLPDLIILPSGDRTEIGKGNLGGQKQKORVSLARAVYSNADIYLFDD  
US-10-665- 733 EEPYRSVIQACALLPDLIILPSGDRTEIGKGNLGGQKQKORVSLARAVYSNADIYLFDD  
US-10-665- 733 EEPYRSVIQACALLPDLIILPSGDRTEIGKGNLGGQKQKORVSLARAVYSNADIYLFDD  
consensus  
US-10-665- 794 PLSAVDAHVGKHI FENVIGPKGMLKNKTRILVTHSMSYLPQVDVII VMSGGKISEMGSYOE  
US-10-665- 794 PLSAVDAHVGKHI FENVIGPKGMLKNKTRILVTHSMSYLPQVDVII VMSGGKISEMGSYOE  
US-10-665- 794 PLSAVDAHVGKHI FENVIGPKGMLKNKTRILVTHSMSYLPQVDVII VMSGGKISEMGSYOE  
US-10-665- 794 PLSAVDAHVGKHI FENVIGPKGMLKNKTRILVTHSMSYLPQVDVII VMSGGKISEMGSYOE  
consensus  
US-10-665- 855 LLARDGAPAEFLRTYASTEQEDAEENGVTGSGPGKEAKQWENGMLVTD SAGKOLORQLS  
US-10-665- 855 LLARDGAPAEFLRTYASTEQEDAEENGVTGSGPGKEAKQWENGMLVTD SAGKOLORQLS  
US-10-665- 855 LLARDGAPAEFLRTYASTEQEDAEENGVTGSGPGKEAKQWENGMLVTD SAGKOLORQLS  
US-10-665- 855 LLARDGAPAEFLRTYASTEQEDAEENGVTGSGPGKEAKQWENGMLVTD SAGKOLORQLS  
consensus  
US-10-665- 916 SSSSYSGDISRHNSHNSAELOKAEAKKEETWKLMEADKAQTOGVKLSVYWDYMKAGLGFISF  
US-10-665- 916 SSSSYSGDISRHNSHNSAELOKAEAKKEETWKLMEADKAQTOGVKLSVYWDYMKAGLGFISF  
US-10-665- 916 SSSSYSGDISRHNSHNSAELOKAEAKKEETWKLMEADKAQTOGVKLSVYWDYMKAGLGFISF  
US-10-665- 916 SSSSYSGDISRHNSHNSAELOKAEAKKEETWKLMEADKAQTOGVKLSVYWDYMKAGLGFISF  
consensus  
US-10-665- 977 LSIPLFMCNHYVSALASNYWLSLWTDPIVNGTQEHKTVRLSVYGALGISQGI AVFGYSMAV  
US-10-665- 977 LSIPLFMCNHYVSALASNYWLSLWTDPIVNGTQEHKTVRLSVYGALGISQGI AVFGYSMAV  
US-10-665- 977 LSIPLFMCNHYVSALASNYWLSLWTDPIVNGTQEHKTVRLSVYGALGISQGI AVFGYSMAV  
US-10-665- 977 LSIPLFMCNHYVSALASNYWLSLWTDPIVNGTQEHKTVRLSVYGALGISQGI AVFGYSMAV  
consensus

US-10-665- 1038 SIGGIIASRCLHVDLLHSILRSPMSFFERTPSGNLVNRF SKELDTVD SMIPEVIKFMGSL  
US-10-665- 1038 SIGGIIASRCLHVDLLHSILRSPMSFFERTPSGNLVNRF SKELDTVD SMIPEVIKFMGSL  
US-10-665- 1038 SIGGIIASRCLHVDLLHSILRSPMSFFERTPSGNLVNRF SKELDTVD SMIPEVIKFMGSL  
US-10-665- 1038 SIGGIIASRCLHVDLLHSILRSPMSFFERTPSGNLVNRF SKELDTVD SMIPEVIKFMGSL  
consensus  
US-10-665- 1099 FNVIGACIVILLATPIAAIIIPPLGLIYFFVQRFYVASSRQLKRLSVSRSPYSHFNETL  
US-10-665- 1099 FNVIGACIVILLATPIAAIIIPPLGLIYFFVQRFYVASSRQLKRLSVSRSPYSHFNETL  
US-10-665- 1099 FNVIGACIVILLATPIAAIIIPPLGLIYFFVQRFYVASSRQLKRLSVSRSPYSHFNETL  
US-10-665- 1099 FNVIGACIVILLATPIAAIIIPPLGLIYFFVQRFYVASSRQLKRLSVSRSPYSHFNETL  
consensus  
US-10-665- 1160 LGVSVIRAFEEQERFIHQSDLKVDENQKAYYPSIVANRWLA VRLECVGNCIVLFAALFAVI  
US-10-665- 1160 LGVSVIRAFEEQERFIHQSDLKVDENQKAYYPSIVANRWLA VRLECVGNCIVLFAALFAVI  
US-10-665- 1160 LGVSVIRAFEEQERFIHQSDLKVDENQKAYYPSIVANRWLA VRLECVGNCIVLFAALFAVI  
US-10-665- 1160 LGVSVIRAFEEQERFIHQSDLKVDENQKAYYPSIVANRWLA VRLECVGNCIVLFAALFAVI  
consensus  
US-10-665- 1221 SRHLSAGLVGLSVSYSLQVTTYLNWLVRMSSEMETNI VAVERLKEYSETEKEAPWQIOET  
US-10-665- 1221 SRHLSAGLVGLSVSYSLQVTTYLNWLVRMSSEMETNI VAVERLKEYSETEKEAPWQIOET  
US-10-665- 1221 SRHLSAGLVGLSVSYSLQVTTYLNWLVRMSSEMETNI VAVERLKEYSETEKEAPWQIOET  
US-10-665- 1221 SRHLSAGLVGLSVSYSLQVTTYLNWLVRMSSEMETNI VAVERLKEYSETEKEAPWQIOET  
consensus  
US-10-665- 1282 APPSSWPQVGRVEFRNYCLRYREDLDFVLRHINVTINGGKVGIVGRTGAGKSSLTGLGFR  
US-10-665- 1282 APPSSWPQVGRVEFRNYCLRYREDLDFVLRHINVTINGGKVGIVGRTGAGKSSLTGLGFR  
US-10-665- 1282 APPSSWPQVGRVEFRNYCLRYREDLDFVLRHINVTINGGKVGIVGRTGAGKSSLTGLGFR  
US-10-665- 1282 APPSSWPQVGRVEFRNYCLRYREDLDFVLRHINVTINGGKVGIVGRTGAGKSSLTGLGFR  
consensus  
US-10-665- 1343 INESAEGEIIIDGINIAKIGLHDLRFKIIIPDPVLFGSLRNMNDPFSQYSDDEEVWTSI  
US-10-665- 1343 INESAEGEIIIDGINIAKIGLHDLRFKIIIPDPVLFGSLRNMNDPFSQYSDDEEVWTSI  
US-10-665- 1343 INESAEGEIIIDGINIAKIGLHDLRFKIIIPDPVLFGSLRNMNDPFSQYSDDEEVWTSI  
US-10-665- 1343 INESAEGEIIIDGINIAKIGLHDLRFKIIIPDPVLFGSLRNMNDPFSQYSDDEEVWTSI  
consensus  
US-10-665- 1404 ELAHLKDFVSALPKLDHECAEGENLSVGORQLVCLARALLRKTILVLDEATAAVDLET  
US-10-665- 1404 ELAHLKDFVSALPKLDHECAEGENLSVGORQLVCLARALLRKTILVLDEATAAVDLET  
US-10-665- 1404 ELAHLKDFVSALPKLDHECAEGENLSVGORQLVCLARALLRKTILVLDEATAAVDLET

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US-10-665- 1404 ELAHLKDFVSALPDKLDHECAGGENLSVGQQLVCLARALKTKIILVDEATAVDLET
consensus
US-10-665- 1465 DDLIQSTIRTPEDCTVLTIAHRLNTIMDYTRVILDKGEIQEYGAPSDLLQORGLFYFMA
US-10-665- 1465 DDLIQSTIRTPEDCTVLTIAHRLNTIMDYTRVILDKGEIQEYGAPSDLLQORGLFYFMA
US-10-665- 1465 DDLIQSTIRTPEDCTVLTIAHRLNTIMDYTRVILDKGEIQEYGAPSDLLQORGLFYFMA
US-10-665- 1465 DDLIQSTIRTPEDCTVLTIAHRLNTIMDYTRVILDKGEIQEYGAPSDLLQORGLFYFMA
consensus
US-10-665- 1526 KDAGLVGGGGGMLSRKGIIPPEEYVLTRLAEDPAEPYRTRERRARFVSKKGCNCNVAHQNI
US-10-665- 1526 KDAGLVGGGGGMLSRKGIIPPEEYVLTRLAEDPAEPYRTRERRARFVSKKGCNCNVAHQNI
US-10-665- 1526 KDAGLVGGGGGMLSRKGIIPPEEYVLTRLAEDPAEPYRTRERRARFVSKKGCNCNVAHQNI
US-10-665- 1526 KDAGLVGGGGGMLSRKGIIPPEEYVLTRLAEDPAEPYRTRERRARFVSKKGCNCNVAHQNI
consensus
US-10-665- 1587 REQGRFLQDVFTTLVDLKWPHHTLLIFTMSFLCSWLLFAMVWMLIAFAHGDIaymekgtd
US-10-665- 1587 REQGRFLQDVFTTLVDLKWPHHTLLIFTMSFLCSWLLFAMVWMLIAFAHGDI
US-10-665- 1587 REQGRFLQDVFTTLVDLKWPHHTLLIFTMSFLCSWLLFAMVWMLIAFAHGDI
US-10-665- 1587 REQGRFLQDVFTTLVDLKWPHHTLLIFTMSFLCSWLLFAMVWMLIAFAHGDI
consensus
US-10-665- 1648 LAPYpydvpdyAGSGTNVPCVTSIHSFSSAFLSIEVQVTIGFGRMVTEECPLAILILIV
US-10-665- 1638 AP GEGTNVPCVTSIHSFSSAFLSIEVQVTIGFGRMVTEECPLAILILIV
US-10-665- 1638 AP GEGTNVPCVTSIHSFSSAFLSIEVQVTIGFGRMVTEECPLAILILIV
US-10-665- 1638 AP GEGTNVPCVTSIHSFSSAFLSIEVQVTIGFGRMVTEECPLAILILIV
consensus
US-10-665- 1709 QNIVGLMINAIMGCI PMKTAQAHRRAETLIFSKHAVITILRHGRLCFMLRVGDLRKSMIIS
US-10-665- 1689 QNIVGLMINAIMGCI PMKTAQAHRRAETLIFSKHAVITILRHGRLCFMLRVGDLRKSMIIS
US-10-665- 1689 QNIVGLMINAIMGCI PMKTAQAHRRAETLIFSKHAVITILRHGRLCFMLRVGDLRKSMIIS
US-10-665- 1689 QNIVGLMINAIMGCI PMKTAQAHRRAETLIFSKHAVITILRHGRLCFMLRVGDLRKSMIIS
consensus
US-10-665- 1770 ATIHMQVVRKTTSPGEVVPVPLHQVDIPMENGVGNGIFLVAPLIIYHVIDSNSPLYDLAPS
US-10-665- 1750 ATIHMQVVRKTTSPGEVVPVPLHQVDIPMENGVGNGIFLVAPLIIYHVIDSNSPLYDLAPS
US-10-665- 1750 ATIHMQVVRKTTSPGEVVPVPLHQVDIPMENGVGNGIFLVAPLIIYHVIDSNSPLYDLAPS
US-10-665- 1750 ATIHMQVVRKTTSPGEVVPVPLHQVDIPMENGVGNGIFLVAPLIIYHVIDSNSPLYDLAPS
consensus
US-10-665- 1831 DLHHQDLEIIVILEGVVETGTTQARTSYLADEILWGQRFVPIVAEEDGRYSVDYSKFG
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US-10-665- 1811 DLHHQDLEIIVILEGVVETGTTQARTSYLADEILWGQRFVPIVAEEDGRYSVDYSKFG
US-10-665- 1811 DLHHQDLEIIVILEGVVETGTTQARTSYLADEILWGQRFVPIVAEEDGRYSVDYSKFG
US-10-665- 1811 DLHHQDLEIIVILEGVVETGTTQARTSYLADEILWGQRFVPIVAEEDGRYSVDYSKFG
consensus
US-10-665- 1892 NTIKVPTPLCTARQLDEDRSLLDALTILASSRGPLRKRSVAVAKAKPKFSISPDLSLS
US-10-665- 1872 NTIKVPTPLCTARQLDEDRSLLDALTILASSRGPLRKRSVAVAKAKPKFSISPDLSLS
US-10-665- 1872 NTIKVPTPLCTARQLDEDRSLLDALTILASSRGPLRaaSVAVAKAKPKFSISPDLSLS
US-10-665- 1872 NTIKVPTPLCTARQLDEDRS
consensus
NTIKVPTPLCTARQLDEDRSLldaltlaasrgplrkrsavakakpkfsisepdals
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Alignment score = 5263.00

Scoring matrix:

|   | 1    | 4    | 6    | 8    |
|---|------|------|------|------|
| 1 |      |      |      |      |
| 4 | 1711 | 1925 | 1827 |      |
| 6 |      | 1711 | 1611 | 1825 |
| 8 |      |      |      |      |

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